



RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/831,631
Source: PUT/09
Date Processed by STIC: 10/29/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

PCT 09

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/831,631

DATE: 10/29/2001
 TIME: 14:16:11

Input Set : A:\6750018999.txt
 Output Set: N:\CRF3\10292001\I831631.raw

Does Not Comply
 Corrected Diskette Needed

3 <110> APPLICANT: Burch, Ronald
 4 Sackler, David
 6 <120> TITLE OF INVENTION: Contraceptive Antibody Vaccines
 8 <130> FILE REFERENCE: 6750-018-999
 10 <140> CURRENT APPLICATION NUMBER: 09/831,631
 11 <141> CURRENT FILING DATE: 2001-09-21
 13 <160> NUMBER OF SEQ ID NOS: 70
 15 <170> SOFTWARE: PatentIn version 3.0
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 16
 19 <212> TYPE: DNA
 C--> 20 <213> ORGANISM: Artificial
 22 <220> FEATURE:
 23 <221> NAME/KEY: misc_feature
 24 <223> OTHER INFORMATION: Description of artificial sequence: Primer for PCR
 26 <400> SEQUENCE: 1
 27 aacagctatg accatg 16
 29 <210> SEQ ID NO: 2
 30 <211> LENGTH: 20
 31 <212> TYPE: DNA
 C--> 32 <213> ORGANISM: Artificial
 34 <220> FEATURE:
 35 <221> NAME/KEY: misc_feature
 36 <223> OTHER INFORMATION: Description of artificial sequence: Primer for PCR
 38 <400> SEQUENCE: 2
 39 gaattcatgg cttgggtgtg 20
 41 <210> SEQ ID NO: 3
 42 <211> LENGTH: 14
 43 <212> TYPE: PRT
 C--> 44 <213> ORGANISM: Artificial
 46 <220> FEATURE:
 47 <221> NAME/KEY: misc_feature
 48 <223> OTHER INFORMATION: Description of artificial sequence: CDR Drived peptide
 50 <220> FEATURE:
 51 <221> NAME/KEY: SITE
 52 <222> LOCATION: (1)..(1)
 53 <223> OTHER INFORMATION: Xaa = biotin Xaa can only represent a single amino acid,
 55 <400> SEQUENCE: 3 nothing else
 W--> 56 Xaa Thr Ala Lys Ala Ser Gln Ser Val Ser Asn Asp Val Ala
 57 1 5 10
 58 <210> SEQ ID NO: 4
 59 <211> LENGTH: 10
 60 <212> TYPE: PRT
 C--> 61 <213> ORGANISM: Artificial
 63 <220> FEATURE:
 64 <221> NAME/KEY: misc_feature
 65 <223> OTHER INFORMATION: Description of artificial sequence: CDR Drived peptide

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67 <220> FEATURE:
68 <221> NAME/KEY: site
69 <222> LOCATION: (1)..(1)
70 <223> OTHER INFORMATION: Xaa = biotin same error
72 <400> SEQUENCE: 4
W--> 73 Xaa Ile Tyr Tyr Ala Ser Asn Arg Tyr Thr
74 1 5 10
75 <210> SEQ ID NO: 5
76 <211> LENGTH: 12
77 <212> TYPE: PRT
C--> 78 <213> ORGANISM: Artificial
80 <220> FEATURE:
81 <221> NAME/KEY: misc_feature
82 <223> OTHER INFORMATION: Description of artificial sequence: CDR Drived peptide
84 <220> FEATURE:
85 <221> NAME/KEY: site
86 <222> LOCATION: (1)..(1)
87 <223> OTHER INFORMATION: Xaa = biotin same
89 <400> SEQUENCE: 5
W--> 90 Xaa Phe Ala Gln Gln Asp Tyr Ser Ser Pro Leu Thr
91 1 5 10
92 <210> SEQ ID NO: 6
93 <211> LENGTH: 8
94 <212> TYPE: PRT
C--> 95 <213> ORGANISM: Artificial
97 <220> FEATURE:
98 <221> NAME/KEY: misc_feature
99 <223> OTHER INFORMATION: Description of artificial sequence: CDR Drived peptide
101 <220> FEATURE:
102 <221> NAME/KEY: site
103 <222> LOCATION: (1)..(1)
104 <223> OTHER INFORMATION: Xaa = biotin same
106 <400> SEQUENCE: 6
W--> 107 Xaa Phe Thr Asn Tyr Gly Met Asn
108 1 5
109 <210> SEQ ID NO: 7
110 <211> LENGTH: 20
111 <212> TYPE: PRT
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114 <220> FEATURE:
115 <221> NAME/KEY: misc_feature
116 <223> OTHER INFORMATION: Description of artificial sequence: CDR Drived peptide
118 <220> FEATURE:
119 <221> NAME/KEY: site
120 <222> LOCATION: (1)..(1)
121 <223> OTHER INFORMATION: Xaa = biotin same
123 <400> SEQUENCE: 7
W--> 124 Xaa Ala Gly Trp Ile Asn Thr Tyr Thr Gly Glu Pro Thr Tyr Ala Asp
125 1 5 10 15

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126 Asp Phe Lys Gly
127      20
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129 <211> LENGTH: 12
130 <212> TYPE: PRT
C--> 131 <213> ORGANISM: Artificial
133 <220> FEATURE:
134 <221> NAME/KEY: misc_feature
135 <223> OTHER INFORMATION: Description of artificial sequence: CDR Drived peptide
137 <220> FEATURE:
138 <221> NAME/KEY: site
139 <222> LOCATION: (1)..(1)
140 <223> OTHER INFORMATION: Xaa = biotin
142 <400> SEQUENCE: 8
W--> 143 Xaa Ala Arg Ala Tyr Tyr Gly Lys Tyr Phe Asp Tyr
144 1      5      10
145 <210> SEQ ID NO: 9
146 <211> LENGTH: 221
147 <212> TYPE: DNA
C--> 148 <213> ORGANISM: Artificial
150 <220> FEATURE:
151 <221> NAME/KEY: misc_feature
152 <223> OTHER INFORMATION: Description of Artificial Sequence: Sperm cell specific
epitope
154 <400> SEQUENCE: 9
155 gaattccagc cttcagggtga acatggctcc ggtgaacagc cttctggtga gcaggcctcg      60
156 ggtgaacagc cttcagggtga gcacgcttca ggggaacagg cttcagggtgc accaatttca      120
157 agcacatcta caggcacaat attaaattgc tacacatgtg cttatatgaa tgatcaagga      180
158 aaatgtcttc gtggagaggg aacctgcatc actcagaatt c      221
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161 <211> LENGTH: 69
162 <212> TYPE: PRT
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165 <220> FEATURE:
166 <221> NAME/KEY: misc_feature
167 <223> OTHER INFORMATION: Description of Artificial Sequence: Sperm cell specific
epitope
169 <400> SEQUENCE: 10
170 Gln Pro Ser Gly Glu His Gly Glu Gln Pro Ser Gly Glu Gln Ala Ser
171 1      5      10      15
172 Gly Glu Gln Pro Ser Gly Glu His Ala Ser Gly Glu Gln Ala Ser Gly
173      20      25      30
174 Ala Gln Ile Ser Ser Thr Ser Thr Gly Thr Ile Leu Asn Cys Tyr Thr
175      35      40      45
176 Cys Ala Tyr Met Asn Asp Gln Gly Lys Cys Leu Arg Gly Glu Gly Thr
177      50      55      60
178 Cys Ile Thr Gln Asn
179 65
180 <210> SEQ ID NO: 11
181 <211> LENGTH: 75
182 <212> TYPE: DNA

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```

C--> 183 <213> ORGANISM: Artificial
      185 <220> FEATURE:
      186 <221> NAME/KEY: misc_feature
      187 <223> OTHER INFORMATION: Description of Artificial Sequence: Cloning primers for SP10
      189 <400> SEQUENCE: 11
      190 gaattccagc cttcagggtga acatgggctcc ggtgaacagc cttctgggtga gcaggcctcg      60
      191 ggtgaacagc cttag                                     75
      193 <210> SEQ ID NO: 12
      194 <211> LENGTH: 75
      195 <212> TYPE: DNA
C--> 196 <213> ORGANISM: Artificial
      198 <220> FEATURE:
      199 <221> NAME/KEY: misc_feature
      200 <223> OTHER INFORMATION: Description of Artificial Sequence: Cloning primers for SP10
      202 <400> SEQUENCE: 12
      203 gtgagcacgc ttcaggggaa cagccttcag gtgcaccaat ttcaagcaca tctacaggca      60
      204 caatattaaa ttgct                                     75
      206 <210> SEQ ID NO: 13
      207 <211> LENGTH: 70
      208 <212> TYPE: DNA
C--> 209 <213> ORGANISM: Artificial
      211 <220> FEATURE:
      212 <221> NAME/KEY: misc_feature
      213 <223> OTHER INFORMATION: Description of Artificial Sequence: Cloning primers for SP10
      215 <400> SEQUENCE: 13
      216 acacatgtgc ttatatgaat gatcaaggaa aatgtcttcg tggagagggga acctgcatca      60
      217 ctcagaattc                                     70
      219 <210> SEQ ID NO: 14
      220 <211> LENGTH: 70
      221 <212> TYPE: DNA
C--> 222 <213> ORGANISM: Artificial
      224 <220> FEATURE:
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      226 <223> OTHER INFORMATION: Description of Artificial Sequence: Cloning primers for SP10
      228 <400> SEQUENCE: 14
      229 acacagcagc ttatatgaat gatcaaggaa aagcacttcg tggagagggga accgcaatca      60
      230 ctcagaattc                                     70
      232 <210> SEQ ID NO: 15
      233 <211> LENGTH: 79
      234 <212> TYPE: DNA
C--> 235 <213> ORGANISM: Artificial
      237 <220> FEATURE:
      238 <221> NAME/KEY: misc_feature
      239 <223> OTHER INFORMATION: Description of Artificial Sequence: Cloning primers for SP10
      241 <400> SEQUENCE: 15
      242 gaattctgag tgatgcagggt tccctctcca cgaagacatt ttccttgatc attcatataa      60
      243 gcacatgtgt agcaattta                                     79
      245 <210> SEQ ID NO: 16
      246 <211> LENGTH: 79

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247 <212> TYPE: DNA
C--> 248 <213> ORGANISM: Artificial
250 <220> FEATURE:
251 <221> NAME/KEY: misc_feature
252 <223> OTHER INFORMATION: Description of Artificial Sequence: Cloning primers for SP10
254 <400> SEQUENCE: 16
255 gaattctgag tgattgcggt tccctctcca cgaagtgcctt tttgatgatc attcatataa      60
256 gctgctgtgt agcaattta                                     79
258 <210> SEQ ID NO: 17
259 <211> LENGTH: 75
260 <212> TYPE: DNA
C--> 261 <213> ORGANISM: Artificial
263 <220> FEATURE:
264 <221> NAME/KEY: misc_feature
265 <223> OTHER INFORMATION: Description of Artificial Sequence: Cloning primers for SP10
267 <400> SEQUENCE: 17
268 atattgtgcc ttagatgtg cttgaaattg gtgcacctga agcctgttcc cctgaagcgt      60
269 gctcacctga aggct                                         75
271 <210> SEQ ID NO: 18
272 <211> LENGTH: 67
273 <212> TYPE: DNA
C--> 274 <213> ORGANISM: Artificial
276 <220> FEATURE:
277 <221> NAME/KEY: misc_feature
278 <223> OTHER INFORMATION: Description of Artificial Sequence: Cloning primers for SP10
280 <400> SEQUENCE: 18
281 gttctcccgga ggcctgctca ccagaaggct gttcacccgga gccatgttca cctgaaggct      60
282 ggaattc                                                  67
284 <210> SEQ ID NO: 19
285 <211> LENGTH: 210
286 <212> TYPE: DNA
C--> 287 <213> ORGANISM: Artificial
289 <220> FEATURE:
290 <221> NAME/KEY: misc_feature
291 <223> OTHER INFORMATION: Description of Artificial Sequence: Sperm cell specific
epitope M
292      SA-6
294 <400> SEQUENCE: 19
295 gtcggcagcc tccgaagcag cccgctccag agcccgtgc tccgaccgct cgtccagagc      60
296 agcctctgct tgetgttctt cttgctgcga tacagctgcg gcgacggcag ctgcagccga      120
297 cgatactgcg acttgacggt gtgccggcga atgtacttgc tgctgcgatt cacggacccg      180
298 ccgctcccgc agacgtgctg cgtcttgagc                          210
300 <210> SEQ ID NO: 20
301 <211> LENGTH: 70
302 <212> TYPE: PRT
C--> 303 <213> ORGANISM: Artificial
305 <220> FEATURE:
306 <221> NAME/KEY: misc_feature
307 <223> OTHER INFORMATION: Description of Artificial Sequence: Sperm cell specific
epitope M
308      SA-6

```

VERIFICATION SUMMARY

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Input Set : A:\6750018999.txt

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L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:20 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:32 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:44 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:56 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:61 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:78 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:95 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:112 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:131 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:148 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:163 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:183 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:196 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:209 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:222 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:235 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:248 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
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L:324 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:21
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L:397 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27
L:409 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:28
L:423 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
L:442 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:30
L:461 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31
L:474 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:32
L:487 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:33
L:500 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:34
L:514 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:35
L:528 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:36
L:542 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:37
L:556 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:38
L:570 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:39
L:584 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:40
L:598 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:41

VERIFICATION SUMMARY

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Input Set : A:\6750018999.txt

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L:612 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:42
L:626 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:43
L:638 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:44
L:651 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:45
L:665 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:46
L:679 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:47
L:692 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:48
L:705 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:49
L:718 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:50